

<110> MOESTRUP, Soren MOLLER, Holger J.

 $<\!120\!>$ THE FUNCTION OF A HAPTOGLOBIN-HAEMOGLOBIN RECEPTOR AND THE USES THEREOF

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<150> DK PA 2001 00039

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<160> 25

<170> PatentIn version 3.1

<210> 1

<211> 347

<212> PRT

<213> Homo sapiens

<400> 1

Met Ser Ala Leu Gly Ala Val Ile Ala Leu Leu Leu Trp Gly Gln Leu 1 5 10 15

Phe Ala Val Asp Ser Gly Asn Asp Val Thr Asp Ile Ala Asp Asp Gly 20 25 30

Cys Pro Lys Pro Pro Glu Ile Ala His Gly Tyr Val Glu His Ser Val 35 40 45

Arg Tyr Gln Cys Lys Asn Tyr Tyr Lys Leu Arg Thr Glu Gly Asp Gly 50 55 60

Val Tyr Thr Leu Asn Asn Glu Lys Gln Trp Ile Asn Lys Ala Val Gly 65 70 75 80

Asp Lys Leu Pro Glu Cys Glu Ala Val Cys Gly Lys Pro Lys Asn Pro 85 90 95

Ala Asn Pro Val Gln Arg Ile Leu Gly Gly His Leu Asp Ala Lys Gly

100 105 110

Ser Phe Pro Trp Gln Ala Lys Met Val Ser His His Asn Leu Thr Thr 115 120 125

- Gly Ala Thr Leu Ile Asn Glu Gln Trp Leu Leu Thr Thr Ala Lys Asn 130 135 140
- Leu Phe Leu Asn His Ser Glu Asn Ala Thr Ala Lys Asp Ile Ala Pro 145 150 155 160
- Thr Leu Thr Leu Tyr Val Gly Lys Lys Gln Leu Val Glu Ile Glu Lys 165 170 175
- Val Val Leu His Pro Asn Tyr Ser Gln Val Asp Ile Gly Leu Ile Lys 180 185 190
- Leu Lys Gln Lys Val Ser Val Asn Glu Arg Val Met Pro Ile Cys Leu 195 200 205
- Pro Ser Lys Asp Tyr Ala Glu Val Gly Arg Val Gly Tyr Val Ser Gly 210 215 220
- Trp Gly Arg Asn Ala Asn Phe Lys Phe Thr Asp His Leu Lys Tyr Val 225 230 230 235
- Met Leu Pro Val Ala Asp Gln Asp Gln Cys Ile Arg His Tyr Glu Gly 245 250 250
- Ser Thr Val Pro Glu Lys Lys Thr Pro Lys Ser Pro Val Gly Val Gln 260 265 270
- Pro Ile Leu Asn Glu His Thr Phe Cys Ala Gly Met Ser Lys Tyr Gln 275 280 285
- Glu Asp Thr Cys Tyr Gly Asp Ala Gly Ser Ala Phe Ala Val His Asp 290 295 300
- Leu Glu Glu Asp Thr Trp Tyr Ala Thr Gly Ile Leu Ser Phe Asp Lys 315 310 315
- Ser Cys Ala Val Ala Glu Tyr Gly Val Tyr Val Lys Val Thr Ser Ile 325 330 335

Gln Asp Trp Val Gln Lys Thr Ile Ala Glu Asn 340 345

<210> 2

<211> 406

<212> PRT

<213> Homo sapiens

<400> 2

Met Ser Ala Leu Gly Ala Val Ile Ala Leu Leu Leu Trp Gly Gln Leu 1 5 10 15

Phe Ala Val Asp Ser Gly Asn Asp Val Thr Asp Ile Ala Asp Asp Gly 20 25 30

Cys Pro Lys Pro Pro Glu Ile Ala His Gly Tyr Val Glu His Ser Val 35 40 45

Arg Tyr Gln Cys Lys Asn Tyr Tyr Lys Leu Arg Thr Glu Gly Asp Gly 50 55 60

Val Tyr Thr Leu Asn Asp Lys Lys Gln Trp Ile Asn Lys Ala Val Gly 65 70 75 80

Asp Lys Leu Pro Glu Cys Glu Ala Asp Asp Gly Cys Pro Lys Pro Pro 85 90 95

Glu Ile Ala His Gly Tyr Val Glu His Ser Val Arg Tyr Gln Cys Lys 100 105 110

Asn Tyr Tyr Lys Leu Arg Thr Glu Gly Asp Gly Val Tyr Thr Leu Asn 115

Asn Glu Lys Gln Trp Ile Asn Lys Ala Val Gly Asp Lys Leu Pro Glu 130 135 140

Cys Glu Ala Val Cys Gly Lys Pro Lys Asn Pro Ala Asn Pro Val Gln 145 150 155 160

Arg Ile Leu Gly Gly His Leu Asp Ala Lys Gly Ser Phe Pro Trp Gln 165 170 175

Ala Lys Met Val Ser His His Asn Leu Thr Thr Gly Ala Thr Leu Ile 180 185 190

Asn Glu Gln Trp Leu Leu Thr Thr Ala Lys Asn Leu Phe Leu Asn His 195 200 205

Ser Glu Asn Ala Thr Ala Lys Asp Ile Ala Pro Thr Leu Thr Leu Tyr 210 215 220

Val Gly Lys Lys Gln Leu Val Glu Ile Glu Lys Val Val Leu His Pro 225 230 235 240

Asn Tyr Ser Gln Val Asp Ile Gly Leu Ile Lys Leu Lys Gln Lys Val 245 250 250

Ser Val Asn Glu Arg Val Met Pro Ile Cys Leu Pro Ser Lys Asp Tyr 260 265 270

Ala Glu Val Gly Arg Val Gly Tyr Val Ser Gly Trp Gly Arg Asn Ala 275 280 285

Asn Phe Lys Phe Thr Asp His Leu Lys Tyr Val Met Leu Pro Val Ala 290 295 300

Asp Gln Asp Gln Cys Ile Arg His Tyr Glu Gly Ser Thr Val Pro Glu 305 310 315 320

Lys Lys Thr Pro Lys Ser Pro Val Gly Val Gln Pro Ile Leu Asn Glu 325 330 335

Gly Asp Ala Gly Ser Ala Phe Ala Val His Asp Leu Glu Glu Asp Thr 355 360 365

Trp Tyr Ala Thr Gly Ile Leu Ser Phe Asp Lys Ser Cys Ala Val Ala 370 375 380

Glu Tyr Gly Val Tyr Val Lys Val Thr Ser Ile Gln Asp Trp Val Gln 385 390 395 400

Lys Thr Ile Ala Glu Asn 405 <210> 3

<211> 347

<212> PRT

<213> Ateles geoffroyi

<400> 3

Met Ser Ala Leu Gly Ala Val Ile Ala Leu Leu Leu Trp Gly Gln Leu 1 5 10 15

Phe Ala Val Asp Ser Gly Asn Asp Val Thr Asp Ile Ala Asp Asp Gly 20 25 30

Cys Pro Lys Pro Pro Glu Ile Ala Asn Gly Tyr Val Glu His Leu Val 35 40 45

Arg Tyr Gln Cys Lys Lys Tyr Tyr Arg Leu Arg Thr Glu Gly Asp Gly 50 55 60

Val Tyr Thr Leu Asn Asn Glu Lys Gln Trp Thr Asn Lys Ala Val Gly 65 70 75 80

Asp Lys Leu Pro Glu Cys Glu Ala Val Cys Gly Lys Pro Lys Asn Pro 85 90 95

Ala Asn Pro Val Gln Arg Ile Leu Gly Gly His Leu Asp Ala Lys Gly
100 105 110

Ser Phe Pro Trp Gln Ala Lys Met Val Ser Arg His Asn Leu Thr Thr 115 120 125

Gly Ala Thr Leu Ile Asn Glu Gln Trp Leu Leu Thr Thr Ala Lys Asn 130 135 140

Leu Phe Leu Asn His Ser Glu Asn Ala Thr Ala Lys Asp Ile Ala Pro 145 150 155 160

Thr Leu Thr Leu Tyr Val Gly Lys Asn Gln Leu Val Glu Ile Glu Lys 165 170 175

Val Val Leu Tyr Pro Asn Tyr Ser Gln Val Asp Ile Gly Leu Ile Lys 180 185 190

Leu Lys Asp Lys Val Pro Val Asn Glu Arg Val Met Pro Ile Cys Leu 195 200 205 Pro Ser Lys Asp Tyr Ala Glu Val Gly Arg Val Gly Tyr Val Ser Gly 210 215 220

Trp Gly Arg Asn Ala Asn Phe Lys Phe Thr Asp His Leu Lys Tyr Val 225 230 235 240

Met Leu Pro Val Ala Asp Gln Tyr Gln Cys Val Lys His Tyr Glu Gly 245 250 255

Ser Thr Val Pro Glu Lys Lys Thr Pro Lys Ser Pro Val Gly Gln Gln 260 265 270

Pro Ile Leu Asn Glu His Thr Phe Cys Ala Gly Met Ser Lys Tyr Gln 275 280 285

Glu Asp Thr Cys Tyr Gly Asp Ala Gly Ser Ala Phe Ala Val His Asp 290 295 300

Leu Glu Glu Asp Thr Trp Tyr Ala Ala Gly Ile Leu Ser Phe Asp Lys 305 310 315 320

Ser Cys Gly Val Ala Glu Tyr Gly Val Tyr Val Lys Ala Thr Ser Ile 325 330 335

Gln Asp Trp Val Gln Lys Thr Ile Ala Glu Asn 340 345

<210> 4

<211> 347

<212> PRT

<213> Mus caroli

<400> 4

Met Arg Ala Leu Gly Ala Val Val Thr Leu Leu Leu Trp Gly Gln Leu 1 5 10 15

Phe Ala Val Glu Leu Gly Asn Asp Ala Met Asp Phe Glu Asp Asp Ser 20 25 30

Cys Pro Lys Pro Pro Glu Ile Ala Asn Gly Tyr Val Glu His Leu Val 35 40 45

Arg Tyr Arg Cys Arg Gln Phe Tyr Arg Leu Arg Ala Glu Gly Asp Gly 50 55 60

Val Tyr Thr Leu Asn Asp Glu Lys Gln Trp Met Asn Thr Val Ala Gly 65 70 75 80

Glu Lys Leu Pro Glu Cys Glu Ala Val Cys Gly Lys Pro Lys His Pro 85 90 95

Val Asp Gln Val Gln Arg Ile Ile Gly Gly Ser Met Asp Ala Lys Gly
100 105 110

Ser Phe Pro Trp Gln Ala Lys Met Ile Ser Arg His Gly Leu Thr Thr 115 120 125

Gly Ala Thr Leu Ile Ser Asp Gln Trp Leu Leu Thr Thr Ala Lys Asn 130 135 140

Leu Phe Leu Asn His Ser Glu Thr Ala Ser Gly Lys Asp Ile Ala Pro 145 150 155 160

Thr Leu Thr Leu Tyr Val Gly Lys Asn Gln Leu Val Glu Ile Glu Lys 165 170 175

Val Ile Leu His Pro Asn His Ser Val Val Asp Ile Gly Leu Ile Lys 180 185 190

Leu Lys Gln Arg Val Leu Val Thr Glu Arg Val Met Pro Ile Cys Leu 195 200 205

Pro Ser Lys Asp Tyr Val Ala Pro Gly Arg Val Gly Tyr Val Ser Gly 210 215 220

Trp Gly Arg Asn Gln Asp Phe Arg Phe Thr Asp Arg Leu Lys Tyr Val 225 230 235 240

Met Leu Pro Val Ala Asp Gln Asp Lys Cys Val Val His Tyr Glu Lys 245 250 255

Ser Thr Val Pro Glu Lys Lys Asn Phe Thr Ser Pro Val Gly Val Gln 260 265 270

Pro Ile Leu Asn Glu His Thr Phe Cys Ala Gly Leu Thr Lys Tyr Glu 275 280 285

Glu Asp Thr Cys Tyr Gly Asp Ala Gly Ser Ala Phe Ala Ile His Asp 290 295

Met Glu Glu Asp Thr Trp Tyr Ala Ala Gly Ile Leu Ser Phe Asp Lys 310

Ser Cys Ala Val Ala Glu Tyr Gly Val Tyr Val Arg Ala Thr Asp Leu 330

Lys Asp Trp Val Gln Glu Thr Met Ala Lys Asn 340

<210> 5

<211> 347 <212> PRT

<213> Mus musculus

<400> 5

Met Arg Ala Leu Gly Ala Val Val Thr Leu Leu Leu Trp Gly Gln Leu

Phe Ala Val Glu Leu Gly Asn Asp Ala Met Asp Phe Glu Asp Asp Ser 25

Cys Pro Lys Pro Pro Glu Ile Ala Asn Gly Tyr Val Glu His Leu Val

Arg Tyr Arg Cys Arg Gln Phe Tyr Arg Leu Arg Ala Glu Gly Asp Gly

Val Tyr Thr Leu Asn Asp Glu Lys Gln Trp Val Asn Thr Val Ala Gly

Glu Lys Leu Pro Glu Cys Glu Ala Val Cys Gly Lys Pro Lys His Pro

Val Asp Gln Val Gln Arg Ile Ile Gly Gly Ser Met Asp Ala Lys Gly

Ser Phe Pro Trp Gln Ala Lys Met Ile Ser Arg His Gly Leu Thr Thr 120

Gly Ala Thr Leu Ile Ser Asp Gln Trp Leu Leu Thr Thr Ala Lys Asn 140

Thr Leu Thr Leu Tyr Val Gly Lys Asn Gln Leu Val Glu Ile Glu Lys 165 170 175

Val Val Leu His Pro Asn His Ser Val Val Asp Ile Gly Leu Ile Lys 180 185 190

Leu Lys Gln Arg Val Leu Val Thr Glu Arg Val Met Pro Ile Cys Leu 195 200 205

Pro Ser Lys Asp Tyr Ile Ala Pro Gly Arg Val Gly Tyr Val Ser Gly 210 215 220

Trp Gly Arg Asn Ala Asn Phe Arg Phe Thr Asp Arg Leu Lys Tyr Val 225 230 235 240

Met Leu Pro Val Ala Asp Gln Asp Lys Cys Val Val His Tyr Glu Asn 245 250 255

Ser Thr Val Pro Glu Lys Lys Asn Leu Thr Ser Pro Val Gly Val Gln 260 265 270

Pro Ile Leu Asn Glu His Thr Phe Cys Ala Gly Leu Thr Lys Tyr Gln 275 280 285

Glu Asp Thr Cys Tyr Gly Asp Ala Gly Ser Ala Phe Ala Ile His Asp 290 295 300

Met Glu Glu Asp Thr Trp Tyr Ala Ala Gly Ile Leu Ser Phe Asp Lys 305 315 315

Ser Cys Ala Val Ala Glu Tyr Gly Val Tyr Val Arg Ala Thr Asp Leu 325 330 335

Lys Asp Trp Val Gln Glu Thr Met Ala Lys Asn 340 345

<210> 6

<211> 347

<212> PRT

<213> Mus saxicola

<220>

<221> misc feature

<222> (311)..(311)

<223> Xaa is unknown

<400> 6

Met Arg Ala Leu Gly Ala Val Val Thr Leu Leu Leu Trp Gly Gln Leu 1 5 10 15

Phe Ala Ala Glu Leu Gly Asn Asp Ala Met Asp Phe Glu Asp Asp Ser 20 25 30

Cys Pro Lys Pro Pro Glu Ile Ala Asn Gly Tyr Val Glu His Leu Val 35 40 45

Arg Tyr Arg Cys Arg Gln Phe Tyr Arg Leu Arg Thr Glu Gly Asp Gly 50 55 60

Val Tyr Thr Leu Asn Asp Glu Lys Gln Trp Val Asn Thr Ala Ala Gly 65 70 . 75 80

Glu Lys Leu Pro Glu Cys Glu Ala Val Cys Gly Lys Pro Lys His Pro 85 90 95

Val Val Gln Val Gln Arg Ile Ile Gly Gly Ser Met Asp Ala Lys Gly
100 105 110

Ser Phe Pro Trp Gln Ala Lys Met Ile Ser Arg His Gly Leu Thr Thr 115 120 125

Gly Ala Thr Leu Ile Ser Asp Gln Trp Leu Leu Thr Thr Ala Lys Asn 130 135 140

Leu Phe Leu Asn His Ser Glu Thr Ala Ser Ala Lys Asp Ile Ala Pro 145 150 155 160

Thr Leu Thr Leu Tyr Val Gly Lys Asn Gln Leu Val Glu Ile Glu Lys 165 170 175

Val Val Leu His Pro Asn His Ser Val Val Asp Ile Gly Leu Ile Lys 180 185 190

Leu Lys Gln Arg Val Leu Val Thr Glu Arg Val Met Pro Ile Cys Leu 195 200 205 Pro Ser Lys Asp Tyr Val Ala Pro Gly Arg Val Gly Tyr Leu Ser Gly 210 215 220

Trp Gly Arg Asn Val Asn Phe Arg Phe Thr Glu Arg Phe Lys Tyr Val 225 230 235 240

Met Leu Pro Val Ala Asp Gln Asp Lys Cys Val Val His Tyr Glu Asn 245 250 255

Ser Thr Val Pro Glu Lys Lys Asn Phe Thr Ser Pro Val Gly Val Gln 260 265 270

Pro Ile Leu Asn Glu His Thr Phe Cys Val Gly Leu Ser Arg Tyr Gln 275 280 285

Glu Asp Thr Cys Tyr Gly Asp Ala Gly Ser Ala Phe Ala Ile His Asp 290 295 300

Met Glu Glu Asp Thr Trp Xaa Ala Ala Gly Ile Leu Ser Phe Asp Lys 305 310 315 320

Ser Cys Ala Val Ala Glu Tyr Gly Val Tyr Val Arg Ala Thr Asp Leu 325 330 335

Lys Asp Trp Val Gln Glu Thr Met Ala Lys Lys 340

<210> 7

<211> 347

<212> PRT

<213> Rattus norvegicus

<400> 7

Met Arg Ala Leu Gly Ala Val Val Thr Leu Leu Leu Trp Gly Gln Leu 1 5 10 15

Phe Ala Val Glu Leu Gly Asn Asp Ala Thr Asp Ile Glu Asp Asp Ser 20 25 30

Cys Pro Lys Pro Pro Glu Ile Ala Asn Gly Tyr Val Glu His Leu Val 35 40 45

Arg Tyr Arg Cys Arg Gln Phe Tyr Lys Leu Gln Thr Glu Gly Asp Gly

50 55 60

Ile Tyr Thr Leu Asn Ser Glu Lys Gln Trp Val Asn Pro Ala Ala Gly 65 70 75 80

Asp Lys Leu Pro Lys Cys Glu Ala Val Cys Gly Lys Pro Lys His Pro 85 90 95

Val Asp Gln Val Gln Arg Ile Ile Gly Gly Ser Met Asp Ala Lys Gly 100 105 110

Ser Phe Pro Trp Gln Ala Lys Met Ile Ser Arg His Gly Leu Thr Thr 115 120 125

Gly Ala Thr Leu Ile Ser Asp Gln Trp Leu Leu Thr Thr Ala Gln Asn 130 135 140

Leu Phe Leu Asn His Ser Glu Asn Ala Thr Ala Lys Asp Ile Ala Pro 145 150 155 160

Thr Leu Thr Leu Tyr Val Gly Lys Asn Gln Leu Val Glu Ile Glu Lys 165 170 175

Val Val Leu His Pro Glu Arg Ser Val Val Asp Ile Gly Leu Ile Lys 180 185 190

Leu Lys Gln Lys Val Leu Val Thr Glu Lys Val Met Pro Ile Cys Leu 195 200 205

Pro Ser Lys Asp Tyr Val Ala Pro Gly Arg Met Gly Tyr Val Ser Gly 210 215 220

Trp Gly Arg Asn Val Asn Phe Arg Phe Thr Glu Arg Leu Lys Tyr Val 225 230 235 240

Met Leu Pro Val Ala Asp Gln Glu Lys Cys Glu Leu His Tyr Glu Lys 245 250 255

Ser Thr Val Pro Glu Lys Lys Gly Ala Val Thr Pro Val Gly Val Gln 260 265 270

Pro Ile Leu Asn Lys His Thr Phe Cys Ala Gly Leu Thr Lys Tyr Glu 275 280 285

Glu Asp Thr Cys Tyr Gly Asp Ala Gly Ser Ala Phe Ala Val His Asp

Thr Glu Glu Asp Thr Trp Tyr Ala Ala Gly Ile Leu Ser Phe Asp Lys

Ser Cys Ala Val Ala Glu Tyr Gly Val Tyr Val Lys Ala Thr Asp Leu 330

Lys Asp Trp Val Gln Glu Thr Met Ala Lys Asn

<210> 8

<211> 346 <212> PRT

<213> Mesocricetus auratus

<400> 8

Met Arg Ala Leu Gly Ala Val Val Thr Leu Leu Leu Trp Gly Gln Leu 1.0

Phe Ala Val Asp Leu Ser Asn Asp Ala Met Asp Thr Ala Asp Asp Ser 20 25 30

Cys Pro Lys Pro Pro Glu Ile Glu Asn Gly Tyr Val Glu His Leu Val

Arg Tyr Arg Cys Gln His Tyr Arg Leu Arg Thr Glu Gly Asp Gly Val 50 55

Tyr Thr Leu Asn Ser Glu Lys Gln Trp Val Asn Thr Ala Ala Gly Glu 70

Arg Leu Pro Glu Cys Glu Ala Val Cys Gly Lys Pro Lys His Pro Val 85 90

Asp Gln Val Gln Arg Ile Ile Gly Gly Ser Leu Asp Ala Lys Gly Ser 100

Phe Pro Trp Gln Ala Lys Met Val Ser Arg His Glu Leu Ile Thr Gly 115 120 125

Ala Thr Leu Ile Ser Asp Gln Trp Leu Leu Thr Thr Ala Lys Asn Leu 130 135

Phe Leu Asn His Ser Glu Asp Ala Thr Ser Lys Asp Ile Ala Pro Thr 145 150 155 160

Leu Lys Leu Tyr Val Gly Lys Met Gln Pro Val Glu Ile Glu Lys Val 165 170 175

Val Ile His Pro Asn Arg Ser Val Val Asp Ile Gly Val Ile Lys Leu 180 185 190

Arg Gln Lys Val Pro Val Asn Glu Arg Val Met Pro Ile Cys Leu Pro 195 200 205

Ser Lys Asp Tyr Ile Ala Pro Gly Arg Met Gly Tyr Val Ser Gly Trp 210 215 220

Gly Arg Asn Ala Asn Phe Arg Phe Thr Asp Arg Leu Lys Tyr Val Met 225 230 235 240

Leu Pro Val Ala Asp Gln Asp Ser Cys Met Leu His Tyr Glu Gly Ser 245 250 255

Thr Val Pro Glu Lys Glu Gly Ser Lys Ser Ser Val Gly Val Gln Pro 260 265 270

Ile Leu Asn Glu His Thr Phe Cys Ala Gly Met Thr Lys Tyr Gln Glu 275 280 285

Asp Thr Cys Tyr Gly Asp Ala Gly Ser Ala Phe Ala Ile His Asp Leu 290 295 300

Glu Gln Asp Thr Trp Tyr Ala Ala Gly Ile Leu Ser Phe Asp Lys Ser 305 310 315 320

Cys Ser Val Ala Glu Tyr Gly Val Tyr Val Lys Val Asn Ser Phe Leu 325 330 335

Asp Trp Ile Gln Glu Thr Met Ala Lys Asn 340 345

<210> 9

<211> 329

<212> PRT

<213> Canis familiaris

<400> 9

Glu Asp Thr Gly Ser Glu Ala Thr Asn Asn Thr Glu Val Ser Leu Pro $1 \hspace{1cm} 5 \hspace{1cm} 10 \hspace{1cm} 15$

Lys Pro Pro Val Ile Glu Asn Gly Tyr Val Glu His Met Ile Arg Tyr 20 25 30

Gln Cys Lys Pro Phe Tyr Lys Leu His Thr Glu Gly Asp Gly Val Tyr 35 40 45

Thr Leu Asn Ser Glu Lys His Trp Thr Asn Lys Ala Val Gly Glu Lys 50 55 60

Leu Pro Glu Cys Glu Ala Val Cys Gly Lys Pro Lys Asn Pro Val Asp 65 70 75 80

Gln Val Gln Arg Ile Met Gly Gly Ser Val Asp Ala Lys Gly Ser Phe 85 90 95

Pro Trp Gln Ala Lys Met Val Ser His His Asn Leu Thr Ser Gly Ala 100 105 110

Thr Leu Ile Asn Glu Gln Trp Leu Leu Thr Thr Ala Lys Asn Leu Phe 115 120 125

Leu Gly His Lys Asp Asp Ala Lys Ala Asn Asp Ile Ala Pro Thr Leu 130 135 140

Lys Leu Tyr Val Gly Lys Asn Gln Leu Val Glu Val Glu Lys Val Val 145 150 155 160

Leu His Pro Asp Tyr Ser Lys Val Asp Ile Gly Leu Ile Lys Leu Lys 165 170 175

Gln Lys Val Pro Ile Asp Glu Arg Val Met Pro Ile Cys Leu Pro Ser 180 185 190

Lys Asp Tyr Ala Glu Val Gly Arg Ile Gly Tyr Val Ser Gly Trp Gly 195 200 205

Arg Asn Ser Asn Phe Asn Phe Thr Glu Leu Leu Lys Tyr Val Met Leu 210 215 220

Pro Val Ala Asp Gln Asp Lys Cys Val Gln His Tyr Glu Gly Ser Thr 225 230 235 240

Val Pro Glu Lys Lys Ser Pro Lys Ser Pro Val Gly Val Gln Pro Ile 245 250 255

Leu Asn Glu His Thr Phe Cys Ala Gly Met Ser Lys Phe Gln Glu Asp 260 265 270

Thr Cys Tyr Gly Asp Ala Gly Ser Ala Phe Ala Val His Asp Gln Asp 275 280 285

Glu Asp Thr Trp Tyr Ala Ala Gly Ile Leu Ser Phe Asp Lys Ser Cys 290 295 300

Thr Val Ala Glu Tyr Gly Val Tyr Val Lys Val Pro Ser Val Leu Ala 305 310 315 320

Trp Val Gln Glu Thr Ile Ala Gly Asn 325

<210> 10

<211> 1116

<212> PRT

<213> Homo sapiens

<400> 10

Met Val Leu Leu Glu Asp Ser Gly Ser Ala Asp Phe Arg Arg His Phe $1 \hspace{1.5cm} 5 \hspace{1.5cm} 10 \hspace{1.5cm} 15$

Val Asn Leu Ser Pro Phe Thr Ile Thr Val Val Leu Leu Ser Ala 20 25 30

Cys Phe Val Thr Ser Ser Leu Gly Gly Thr Asp Lys Glu Leu Arg Leu 35 40 45

Val Asp Gly Glu Asn Lys Cys Ser Gly Arg Val Glu Val Lys Val Gln 50 55 60

Glu Glu Trp Gly Thr Val Cys Asn Asn Gly Trp Ser Met Glu Ala Val 65 70 75 80

Ser Val Ile Cys Asn Gln Leu Gly Cys Pro Thr Ala Ile Lys Ala Pro 85 90 95

- Gly Trp Ala Asn Ser Ser Ala Gly Ser Gly Arg Ile Trp Met Asp His 100 105 110
- Val Ser Cys Arg Gly Asn Glu Ser Ala Leu Trp Asp Cys Lys His Asp 115 120 125
- Gly Trp Gly Lys His Ser Asn Cys Thr His Gln Gln Asp Ala Gly Val 130 135 140
- Thr Cys Ser Asp Gly Ser Asn Leu Glu Met Arg Leu Thr Arg Gly Gly 145 150 155 160
- Asn Met Cys Ser Gly Arg Ile Glu Ile Lys Phe Gln Gly Arg Trp Gly 165 170 175
- Thr Val Cys Asp Asp Asn Phe Asn Ile Asp His Ala Ser Val Ile Cys
 180 . 185 . 190
- Arg Gln Leu Glu Cys Gly Ser Ala Val Ser Phe Ser Gly Ser Ser Asn 195 200 205
- Phe Gly Glu Gly Ser Gly Pro Ile Trp Phe Asp Asp Leu Ile Cys Asn 210 215 220
- Gly Asn Glu Ser Ala Leu Trp Asn Cys Lys His Gln Gly Trp Gly Lys 225 230 235 240
- His Asn Cys Asp His Ala Glu Asp Ala Gly Val Ile Cys Ser Lys Gly 245 250 255
- Ala Asp Leu Ser Leu Arg Leu Val Asp Gly Val Thr Glu Cys Ser Gly 260 265 270
- Arg Leu Glu Val Arg Phe Gln Gly Glu Trp Gly Thr Ile Cys Asp Asp 275 280 285
- Gly Trp Asp Ser Tyr Asp Ala Ala Val Ala Cys Lys Gln Leu Gly Cys 290 295 300
- Pro Thr Ala Val Thr Ala Ile Gly Arg Val Asn Ala Ser Lys Gly Phe 305 310 315 320

Gly His Ile Trp Leu Asp Ser Val Ser Cys Gln Gly His Glu Pro Ala 325 330 335

Val Trp Gln Cys Lys His His Glu Trp Gly Lys His Tyr Cys Asn His 340 345 350

Asn Glu Asp Ala Gly Val Thr Cys Ser Asp Gly Ser Asp Leu Glu Leu 355 360 365

Arg Leu Arg Gly Gly Gly Ser Arg Cys Ala Gly Thr Val Glu Val Glu 370 380

Ile Gln Arg Leu Leu Gly Lys Val Cys Asp Arg Gly Trp Gly Leu Lys 385 390 395 400

Glu Ala Asp Val Val Cys Arg Gln Leu Gly Cys Gly Ser Ala Leu Lys $405 \hspace{1.5cm} 410 \hspace{1.5cm} 415 \hspace{1.5cm}$

Thr Ser Tyr Gln Val Tyr Ser Lys Ile Gln Ala Thr Asn Thr Trp Leu 420 425 430

Phe Leu Ser Ser Cys Asn Gly Asn Glu Thr Ser Leu Trp Asp Cys Lys 435 440 445

Asn Trp Gln Trp Gly Gly Leu Thr Cys Asp His Tyr Glu Glu Ala Lys 450 455 460

Ile Thr Cys Ser Ala His Arg Glu Pro Arg Leu Val Gly Gly Asp Ile 465 470 475 480

Pro Cys Ser Gly Arg Val Glu Val Lys His Gly Asp Thr Trp Gly Ser 485 490 495

Ile Cys Asp Ser Asp Phe Ser Leu Glu Ala Ala Ser Val Leu Cys Arg 500 505 510

Glu Leu Gln Cys Gly Thr Val Val Ser Ile Leu Gly Gly Ala His Phe 515 520 525

Gly Glu Gly Asn Gly Gln Ile Trp Ala Glu Glu Phe Gln Cys Glu Gly 530 535 540

His Glu Ser His Leu Ser Leu Cys Pro Val Ala Pro Arg Pro Glu Gly 545 550 555 560

Thr Cys Ser His Ser Arg Asp Val Gly Val Val Cys Ser Arg Tyr Thr 565 570 575

Glu Ile Arg Leu Val Asn Gly Lys Thr Pro Cys Glu Gly Arg Val Glu 580 585 590

Leu Lys Thr Leu Gly Ala Trp Gly Ser Leu Cys Asn Ser His Trp Asp 595 600 605

Ile Glu Asp Ala His Val Leu Cys Gln Gln Leu Lys Cys Gly Val Ala 610 615 620

Leu Ser Thr Pro Gly Gly Ala Arg Phe Gly Lys Gly Asn Gly Gln Ile 625 630 635 640

Trp Arg His Met Phe His Cys Thr Gly Thr Glu Gln His Met Gly Asp
645 650 655

Cys Pro Val Thr Ala Leu Gly Ala Ser Leu Cys Pro Ser Glu Gln Val 660 665 670

Ala Ser Val Ile Cys Ser Gly Asn Gln Ser Gln Thr Leu Ser Ser Cys 675 680 685

Asn Ser Ser Ser Leu Gly Pro Thr Arg Pro Thr Ile Pro Glu Glu Ser 690 695 700

Ala Val Ala Cys Ile Glu Ser Gly Gln Leu Arg Leu Val Asn Gly Gly 705 710 715 720

Gly Arg Cys Ala Gly Arg Val Glu Ile Tyr His Glu Gly Ser Trp Gly
725 730 735

Thr Ile Cys Asp Asp Ser Trp Asp Leu Ser Asp Ala His Val Val Cys
740 745 750

Arg Gln Leu Gly Cys Gly Glu Ala Ile Asn Ala Thr Gly Ser Ala His 755 760 765

Phe Gly Glu Gly Thr Gly Pro Ile Trp Leu Asp Glu Met Lys Cys Asn 770 780

Gly Lys Glu Ser Arg Ile Trp Gln Cys His Ser His Gly Trp Gly Gln 785 790 795 800

Gln Asn Cys Arg His Lys Glu Asp Ala Gly Val Ile Cys Ser Glu Phe 805 810 815

Met Ser Leu Arg Leu Thr Ser Glu Ala Ser Arg Glu Ala Cys Ala Gly 820 825 830

Arg Leu Glu Val Phe Tyr Asn Gly Ala Trp Gly Thr Val Gly Lys Ser 835 840 845

Ser Met Ser Glu Thr Thr Val Gly Val Val Cys Arg Gln Leu Gly Cys 850 855 860

Ala Asp Lys Gly Lys Ile Asn Pro Ala Ser Leu Asp Lys Ala Met Ser 865 870 875 880

Ile Pro Met Trp Val Asp Asn Val Gln Cys Pro Lys Gly Pro Asp Thr 885 890 895

Leu Trp Gln Cys Pro Ser Ser Pro Trp Glu Lys Arg Leu Ala Ser Pro 900 905 910

Ser Glu Glu Thr Trp Ile Thr Cys Asp Asn Lys Ile Arg Leu Gln Glu 915 920 925

Gly Pro Thr Ser Cys Ser Gly Arg Val Glu Ile Trp His Gly Gly Ser 930 935 940

Trp Gly Thr Val Cys Asp Asp Ser Trp Asp Leu Asp Asp Ala Gln Val 945 950 955 960

Val Cys Gln Gln Leu Gly Cys Gly Pro Ala Leu Lys Ala Phe Lys Glu 965 970 975

Ala Glu Phe Gly Gln Gly Thr Gly Pro Ile Trp Leu Asn Glu Val Lys 980 985 990

Cys Lys Gly Asn Glu Ser Ser Leu Trp Asp Cys Pro Ala Arg Arg Trp 995 1000 1005

Gly His Ser Glu Cys Gly His Lys Glu Asp Ala Ala Val Asn Cys 1010 1015 1020 Thr Asp Ile Ser Val Gln Lys Thr Pro Gln Lys Ala Thr Thr Gly 1025 1030

Arg Ser Ser Arg Gln Ser Ser Phe Ile Ala Val Gly Ile Leu Gly

Val Val Leu Leu Ala Ile Phe Val Ala Leu Phe Phe Leu Thr Lys 1055

Lys Arg Arg Gln Arg Gln Arg Leu Ala Val Ser Ser Arg Gly Glu 1070 1075

Asn Leu Val His Gln Ile Gln Tyr Arg Glu Met Asn Ser Cys Leu 1085 1090

Asn Ala Asp Asp Leu Asp Leu Met Asn Ser Ser Gly Gly His Ser 1100 1105

Glu Pro His 1115

<210> 11 <211> 1149 <212> PRT

<213> Homo sapiens

<400> 11

Met Val Leu Leu Glu Asp Ser Gly Ser Ala Asp Phe Arg Arg His Phe 10

Val Asn Leu Ser Pro Phe Thr Ile Thr Val Val Leu Leu Ser Ala 20 . 25

Cys Phe Val Thr Ser Ser Leu Gly Gly Thr Asp Lys Glu Leu Arg Leu

Val Asp Gly Glu Asn Lys Cys Ser Gly Arg Val Glu Val Lys Val Gln

Glu Glu Trp Gly Thr Val Cys Asn Asn Gly Trp Ser Met Glu Ala Val

Ser Val Ile Cys Asn Gln Leu Gly Cys Pro Thr Ala Ile Lys Ala Pro

90 95

Gly Trp Ala Asn Ser Ser Ala Gly Ser Gly Arg Ile Trp Met Asp His 100 105 110

Val Ser Cys Arg Gly Asn Glu Ser Ala Leu Trp Asp Cys Lys His Asp 115 120 125

Gly Trp Gly Lys His Ser Asn Cys Thr His Gln Gln Asp Ala Gly Val 130 135

Thr Cys Ser Asp Gly Ser Asn Leu Glu Met Arg Leu Thr Arg Gly Gly 145 150 155 160

Asn Met Cys Ser Gly Arg Ile Glu Ile Lys Phe Gln Gly Arg Trp Gly
165 170 175

Thr Val Cys Asp Asp Asn Phe Asn Ile Asp His Ala Ser Val Ile Cys 180 185 190

Arg Gln Leu Glu Cys Gly Ser Ala Val Ser Phe Ser Gly Ser Ser Asn 195 200 205

Phe Gly Glu Gly Ser Gly Pro Ile Trp Phe Asp Asp Leu Ile Cys Asn 210 220

Gly Asn Glu Ser Ala Leu Trp Asn Cys Lys His Gln Gly Trp Gly Lys 225 230 235 240

His Asn Cys Asp His Ala Glu Asp Ala Gly Val Ile Cys Ser Lys Gly 245 250 255

Ala Asp Leu Ser Leu Arg Leu Val Asp Gly Val Thr Glu Cys Ser Gly 260 265 270

Arg Leu Glu Val Arg Phe Gln Gly Glu Trp Gly Thr Ile Cys Asp Asp 275 280 285

Gly Trp Asp Ser Tyr Asp Ala Ala Val Ala Cys Lys Gln Leu Gly Cys 290 295 300

Pro Thr Ala Val Thr Ala Ile Gly Arg Val Asn Ala Ser Lys Gly Phe 305 310 315 320 Gly His Ile Trp Leu Asp Ser Val Ser Cys Gln Gly His Glu Pro Ala 325 330 335

Val Trp Gln Cys Lys His His Glu Trp Gly Lys His Tyr Cys Asn His 340 345 350

Asn Glu Asp Ala Gly Val Thr Cys Ser Asp Gly Ser Asp Leu Glu Leu 355 360 365

Arg Leu Arg Gly Gly Gly Ser Arg Cys Ala Gly Thr Val Glu Val Glu 370 375 380

Ile Gln Arg Leu Leu Gly Lys Val Cys Asp Arg Gly Trp Gly Leu Lys 385 390 395 400

Glu Ala Asp Val Val Cys Arg Gln Leu Gly Cys Gly Ser Ala Leu Lys 405 410 415

Thr Ser Tyr Gln Val Tyr Ser Lys Ile Gln Ala Thr Asn Thr Trp Leu 420 425 430

Phe Leu Ser Ser Cys Asn Gly Asn Glu Thr Ser Leu Trp Asp Cys Lys 435

Asn Trp Gln Trp Gly Gly Leu Thr Cys Asp His Tyr Glu Glu Ala Lys 450 455 460

Ile Thr Cys Ser Ala His Arg Glu Pro Arg Leu Val Gly Gly Asp Ile465470475480

Pro Cys Ser Gly Arg Val Glu Val Lys His Gly Asp Thr Trp Gly Ser 485 490 495

Ile Cys Asp Ser Asp Phe Ser Leu Glu Ala Ala Ser Val Leu Cys Arg 500 505 510

Glu Leu Gln Cys Gly Thr Val Val Ser Ile Leu Gly Gly Ala His Phe 515 520 525

Gly Glu Gly Asn Gly Gln Ile Trp Ala Glu Glu Phe Gln Cys Glu Gly 530 540

His Glu Ser His Leu Ser Leu Cys Pro Val Ala Pro Arg Pro Glu Gly

545 550 555 560 Thr Cys Ser His Ser Arg Asp Val Gly Val Val Cys Ser Ser Lys Thr 565 570 Gln Lys Thr Ser Leu Ile Gly Ser Tyr Thr Val Lys Gly Thr Gly Leu 585 Gly Ser His Ser Cys Leu Phe Leu Lys Pro Cys Leu Leu Pro Gly Tyr Thr Glu Ile Arg Leu Val Asn Gly Lys Thr Pro Cys Glu Gly Arg Val 610 Glu Leu Lys Thr Leu Gly Ala Trp Gly Ser Leu Cys Asn Ser His Trp 630 Asp Ile Glu Asp Ala His Val Leu Cys Gln Gln Leu Lys Cys Gly Val 645 650 Ala Leu Ser Thr Pro Gly Gly Ala Arg Phe Gly Lys Gly Asn Gly Gln 665 Ile Trp Arg His Met Phe His Cys Thr Gly Thr Glu Gln His Met Gly 675 Asp Cys Pro Val Thr Ala Leu Gly Ala Ser Leu Cys Pro Ser Glu Gln 695 Val Ala Ser Val Ile Cys Ser Gly Asn Gln Ser Gln Thr Leu Ser Ser Cys Asn Ser Ser Leu Gly Pro Thr Arg Pro Thr Ile Pro Glu Glu 730 Ser Ala Val Ala Cys Ile Glu Ser Gly Gln Leu Arg Leu Val Asn Gly Gly Gly Arg Cys Ala Gly Arg Val Glu Ile Tyr His Glu Gly Ser Trp Gly Thr Ile Cys Asp Asp Ser Trp Asp Leu Ser Asp Ala His Val Val

770

Cys Arg Gln Leu Gly Cys Gly Glu Ala Ile Asn Ala Thr Gly Ser Ala 785 790 795 800

His Phe Gly Glu Gly Thr Gly Pro Ile Trp Leu Asp Glu Met Lys Cys 805 810 815

Asn Gly Lys Glu Ser Arg Ile Trp Gln Cys His Ser His Gly Trp Gly 820 825 830

Gln Gln Asn Cys Arg His Lys Glu Asp Ala Gly Val Ile Cys Ser Glu 835 840 845

Phe Met Ser Leu Arg Leu Thr Ser Glu Ala Ser Arg Glu Ala Cys Ala 850 855 860

Gly Arg Leu Glu Val Phe Tyr Asn Gly Ala Trp Gly Thr Val Gly Lys 865 870 875 880

Ser Ser Met Ser Glu Thr Thr Val Gly Val Val Cys Arg Gln Leu Gly 885 890 895

Cys Ala Asp Lys Gly Lys Ile Asn Pro Ala Ser Leu Asp Lys Ala Met 900 905 910

Ser Ile Pro Met Trp Val Asp Asn Val Gln Cys Pro Lys Gly Pro Asp 915 920 925

Thr Leu Trp Gln Cys Pro Ser Ser Pro Trp Glu Lys Arg Leu Ala Ser 930 935 940

Pro Ser Glu Glu Thr Trp Ile Thr Cys Asp Asn Lys Ile Arg Leu Gln 945 950 955 960

Glu Gly Pro Thr Ser Cys Ser Gly Arg Val Glu Ile Trp His Gly Gly
965 970 975

Ser Trp Gly Thr Val Cys Asp Asp Ser Trp Asp Leu Asp Asp Ala Gln 980 985 990

Val Val Cys Gln Gln Leu Gly Cys Gly Pro Ala Leu Lys Ala Phe Lys 995 1000 1005

Glu Ala Glu Phe Gly Gln Gly Thr Gly Pro Ile Trp Leu Asn Glu

1010 1015 1020 Val Lys Cys Lys Gly Asn Glu Ser Ser Leu Trp Asp Cys Pro Ala 1030 1025 1035 Arg Arg Trp Gly His Ser Glu Cys Gly His Lys Glu Asp Ala Ala 1045 1050 Val Asn Cys Thr Asp Ile Ser Val Gln Lys Thr Pro Gln Lys Ala 1060 Thr Thr Gly Arg Ser Ser Arg Gln Ser Ser Phe Ile Ala Val Gly 1070 1075 Ile Leu Gly Val Val Leu Leu Ala Ile Phe Val Ala Leu Phe Phe 1085 1090 1095 Leu Thr Lys Lys Arg Arg Gln Arg Gln Arg Leu Ala Val Ser Ser 1100 1105 1110 Arg Gly Glu Asn Leu Val His Gln Ile Gln Tyr Arg Glu Met Asn 1115 1120 Ser Cys Leu Asn Ala Asp Asp Leu Asp Leu Met Asn Ser Ser Gly 1130 1135 Gly His Ser Glu Pro His 1145 <210> 12 <211> 1156 <212> PRT <213> Homo sapiens <400> 12 Met Val Leu Leu Glu Asp Ser Gly Ser Ala Asp Phe Arg Arg His Phe

Cys Phe Val Thr Ser Ser Leu Gly Gly Thr Asp Lys Glu Leu Arg Leu 35 40 45

Val Asn Leu Ser Pro Phe Thr Ile Thr Val Val Leu Leu Ser Ala

25

10

5

20 .

Val Asp Gly Glu Asn Lys Cys Ser Gly Arg Val Glu Val Lys Val Gln 50 55 60

Glu Glu Trp Gly Thr Val Cys Asn Asn Gly Trp Ser Met Glu Ala Val 65 70 75 80

Ser Val Ile Cys Asn Gln Leu Gly Cys Pro Thr Ala Ile Lys Ala Pro 85 90 95

Gly Trp Ala Asn Ser Ser Ala Gly Ser Gly Arg Ile Trp Met Asp His 100 105 110

Val Ser Cys Arg Gly Asn Glu Ser Ala Leu Trp Asp Cys Lys His Asp 115 120 125

Gly Trp Gly Lys His Ser Asn Cys Thr His Gln Gln Asp Ala Gly Val 130 135 140

Thr Cys Ser Asp Gly Ser Asn Leu Glu Met Arg Leu Thr Arg Gly Gly 145 150 155 160

Asn Met Cys Ser Gly Arg Ile Glu Ile Lys Phe Gln Gly Arg Trp Gly 165 170 175

Thr Val Cys Asp Asp Asn Phe Asn Ile Asp His Ala Ser Val Ile Cys
180 185 190

Arg Gln Leu Glu Cys Gly Ser Ala Val Ser Phe Ser Gly Ser Ser Asn 195 200 205

Phe Gly Glu Gly Ser Gly Pro Ile Trp Phe Asp Asp Leu Ile Cys Asn 210 225

Gly Asn Glu Ser Ala Leu Trp Asn Cys Lys His Gln Gly Trp Gly Lys 225 230 235 240

His Asn Cys Asp His Ala Glu Asp Ala Gly Val Ile Cys Ser Lys Gly 245 250 255

Ala Asp Leu Ser Leu Arg Leu Val Asp Gly Val Thr Glu Cys Ser Gly 260 265 270

Arg Leu Glu Val Arg Phe Gln Gly Glu Trp Gly Thr Ile Cys Asp Asp 275 280 285

- Gly Trp Asp Ser Tyr Asp Ala Ala Val Ala Cys Lys Gln Leu Gly Cys 290 295 300
- Pro Thr Ala Val Thr Ala Ile Gly Arg Val Asn Ala Ser Lys Gly Phe 305 310 315 320
- Gly His Ile Trp Leu Asp Ser Val Ser Cys Gln Gly His Glu Pro Ala 325 330 335
- Val Trp Gln Cys Lys His His Glu Trp Gly Lys His Tyr Cys Asn His 340 345 350
- Asn Glu Asp Ala Gly Val Thr Cys Ser Asp Gly Ser Asp Leu Glu Leu 355 360 365
- Arg Leu Arg Gly Gly Ser Arg Cys Ala Gly Thr Val Glu Val Glu 370 375 380
- Ile Gln Arg Leu Leu Gly Lys Val Cys Asp Arg Gly Trp Gly Leu Lys 385 390 395 400
- Glu Ala Asp Val Val Cys Arg Gln Leu Gly Cys Gly Ser Ala Leu Lys 405 410 415
- Thr Ser Tyr Gln Val Tyr Ser Lys Ile Gln Ala Thr Asn Thr Trp Leu 420 425 430
- Phe Leu Ser Ser Cys Asn Gly Asn Glu Thr Ser Leu Trp Asp Cys Lys 435 440 445
- Asn Trp Gln Trp Gly Gly Leu Thr Cys Asp His Tyr Glu Glu Ala Lys 450 460
- Ile Thr Cys Ser Ala His Arg Glu Pro Arg Leu Val Gly Gly Asp Ile 465 470 475 480
- Pro Cys Ser Gly Arg Val Glu Val Lys His Gly Asp Thr Trp Gly Ser 485 490 495
- Ile Cys Asp Ser Asp Phe Ser Leu Glu Ala Ala Ser Val Leu Cys Arg 500 505 510

- Glu Leu Gln Cys Gly Thr Val Val Ser Ile Leu Gly Gly Ala His Phe 515 525
- Gly Glu Gly Asn Gly Gln Ile Trp Ala Glu Glu Phe Gln Cys Glu Gly 530 535 540
- His Glu Ser His Leu Ser Leu Cys Pro Val Ala Pro Arg Pro Glu Gly 545 550 555 560
- Thr Cys Ser His Ser Arg Asp Val Gly Val Val Cys Ser Arg Tyr Thr 565 570 575
- Glu Ile Arg Leu Val Asn Gly Lys Thr Pro Cys Glu Gly Arg Val Glu 580 585 590
- Leu Lys Thr Leu Gly Ala Trp Gly Ser Leu Cys Asn Ser His Trp Asp 595 600 605
- Ile Glu Asp Ala His Val Leu Cys Gln Gln Leu Lys Cys Gly Val Ala 610 615 620
- Leu Ser Thr Pro Gly Gly Ala Arg Phe Gly Lys Gly Asn Gly Gln Ile 625 630 635 635
- Trp Arg His Met Phe His Cys Thr Gly Thr Glu Gln His Met Gly Asp 645 650 655
- Cys Pro Val Thr Ala Leu Gly Ala Ser Leu Cys Pro Ser Glu Gln Val 660 665 670
- Ala Ser Val Ile Cys Ser Gly Asn Gln Ser Gln Thr Leu Ser Ser Cys 675 680 685
- Asn Ser Ser Ser Leu Gly Pro Thr Arg Pro Thr Ile Pro Glu Glu Ser 690 695 700
- Ala Val Ala Cys Ile Glu Ser Gly Gln Leu Arg Leu Val Asn Gly Gly 705 710 715 720
- Gly Arg Cys Ala Gly Arg Val Glu Ile Tyr His Glu Gly Ser Trp Gly
 725 730 735
- Thr Ile Cys Asp Asp Ser Trp Asp Leu Ser Asp Ala His Val Val Cys 740 745 750

Arg Gln Leu Gly Cys Gly Glu Ala Ile Asn Ala Thr Gly Ser Ala His 755 760 765

Phe Gly Glu Gly Thr Gly Pro Ile Trp Leu Asp Glu Met Lys Cys Asn 770 775 780

Gly Lys Glu Ser Arg Ile Trp Gln Cys His Ser His Gly Trp Gly Gln
785 790 795 800

Gln Asn Cys Arg His Lys Glu Asp Ala Gly Val Ile Cys Ser Glu Phe 805 810 815

Met Ser Leu Arg Leu Thr Ser Glu Ala Ser Arg Glu Ala Cys Ala Gly 820 825 830

Arg Leu Glu Val Phe Tyr Asn Gly Ala Trp Gly Thr Val Gly Lys Ser 835 840 845

Ser Met Ser Glu Thr Thr Val Gly Val Val Cys Arg Gln Leu Gly Cys 850 855 860

Ala Asp Lys Gly Lys Ile Asn Pro Ala Ser Leu Asp Lys Ala Met Ser 865 870 875 880

Ile Pro Met Trp Val Asp Asn Val Gln Cys Pro Lys Gly Pro Asp Thr 885 890 895

Leu Trp Gln Cys Pro Ser Ser Pro Trp Glu Lys Arg Leu Ala Ser Pro 900 905 910

Ser Glu Glu Thr Trp Ile Thr Cys Asp Asn Lys Ile Arg Leu Gln Glu 915 920 925

Gly Pro Thr Ser Cys Ser Gly Arg Val Glu Ile Trp His Gly Gly Ser 930 935 940

Trp Gly Thr Val Cys Asp Asp Ser Trp Asp Leu Asp Asp Ala Gln Val 945 950 955 960

Val Cys Gln Gln Leu Gly Cys Gly Pro Ala Leu Lys Ala Phe Lys Glu 965 970 975 Ala Glu Phe Gly Gln Gly Thr Gly Pro Ile Trp Leu Asn Glu Val Lys 980

Cys Lys Gly Asn Glu Ser Ser Leu Trp Asp Cys Pro Ala Arg Arg Trp 995 1000 1005

Gly His Ser Glu Cys Gly His Lys Glu Asp Ala Ala Val Asn Cys 1015 1010

Thr Asp Ile Ser Val Gln Lys Thr Pro Gln Lys Ala Thr Thr Gly 1025 1030

Arg Ser Ser Arg Gln Ser Ser Phe Ile Ala Val Gly Ile Leu Gly 1040 1045

Val Val Leu Leu Ala Ile Phe Val Ala Leu Phe Phe Leu Thr Lys 1055 1060 1065

Lys Arg Arg Gln Arg Gln Arg Leu Ala Val Ser Ser Arg Gly Glu 1070 1075 1080

Asn Leu Val His Gln Ile Gln Tyr Arg Glu Met Asn Ser Cys Leu 1085 1090 1095

Asn Ala Asp Asp Leu Asp Leu Met Asn Ser Ser Gly Leu Trp Val 1100 1105 1110

Leu Gly Gly Ser Ile Ala Gln Gly Phe Arg Ser Val Ala Ala Val 1115 1120

Glu Ala Gln Thr Phe Tyr Phe Asp Lys Gln Leu Lys Lys Ser Lys 1130 1135 1140

Asn Val Ile Gly Ser Leu Asp Ala Tyr Asn Gly Gln Glu 1150 1155

<210> 13 <211> 1151 <212> PRT <213> Homo sapiens

<400> 13

Met Val Leu Leu Glu Asp Ser Gly Ser Ala Asp Phe Arg Arg His Phe

Val Asn Leu Ser Pro Phe Thr Ile Thr Val Val Leu Leu Ser Ala 20 25 30

Cys Phe Val Thr Ser Ser Leu Gly Gly Thr Asp Lys Glu Leu Arg Leu 35 40 45

Val Asp Gly Glu Asn Lys Cys Ser Gly Arg Val Glu Val Lys Val Gln 50 55 60

Glu Glu Trp Gly Thr Val Cys Asn Asn Gly Trp Ser Met Glu Ala Val 65 70 75 80

Ser Val Ile Cys Asn Gln Leu Gly Cys Pro Thr Ala Ile Lys Ala Pro 85 90 95

Gly Trp Ala Asn Ser Ser Ala Gly Ser Gly Arg Ile Trp Met Asp His 100 105 110

Val Ser Cys Arg Gly Asn Glu Ser Ala Leu Trp Asp Cys Lys His Asp 115 120 125

Gly Trp Gly Lys His Ser Asn Cys Thr His Gln Gln Asp Ala Gly Val 130 135

Thr Cys Ser Asp Gly Ser Asn Leu Glu Met Arg Leu Thr Arg Gly Gly 145 150 155 160

Asn Met Cys Ser Gly Arg Ile Glu Ile Lys Phe Gln Gly Arg Trp Gly 165 170 175

Thr Val Cys Asp Asp Asn Phe Asn Ile Asp His Ala Ser Val Ile Cys 180 185 190

Arg Gln Leu Glu Cys Gly Ser Ala Val Ser Phe Ser Gly Ser Ser Asn 195 200 205

Phe Gly Glu Gly Ser Gly Pro Ile Trp Phe Asp Asp Leu Ile Cys Asn 210 215 220

Gly Asn Glu Ser Ala Leu Trp Asn Cys Lys His Gln Gly Trp Gly Lys 225 230 235 240

His Asn Cys Asp His Ala Glu Asp Ala Gly Val Ile Cys Ser Lys Gly

245 250 255

Ala Asp Leu Ser Leu Arg Leu Val Asp Gly Val Thr Glu Cys Ser Gly 260 265 270

Arg Leu Glu Val Arg Phe Gln Gly Glu Trp Gly Thr Ile Cys Asp Asp 275 280 285

Gly Trp Asp Ser Tyr Asp Ala Ala Val Ala Cys Lys Gln Leu Gly Cys 290 295 300

Pro Thr Ala Val Thr Ala Ile Gly Arg Val Asn Ala Ser Lys Gly Phe 305 310 315

Gly His Ile Trp Leu Asp Ser Val Ser Cys Gln Gly His Glu Pro Ala 325 330 335

Val Trp Gln Cys Lys His His Glu Trp Gly Lys His Tyr Cys Asn His 340 345 350

Asn Glu Asp Ala Gly Val Thr Cys Ser Asp Gly Ser Asp Leu Glu Leu 355 360 365

Arg Leu Arg Gly Gly Ser Arg Cys Ala Gly Thr Val Glu Val Glu 370 375 380

Ile Gln Arg Leu Leu Gly Lys Val Cys Asp Arg Gly Trp Gly Leu Lys 385 390 395 400

Glu Ala Asp Val Val Cys Arg Gln Leu Gly Cys Gly Ser Ala Leu Lys 405 410 415

Thr Ser Tyr Gln Val Tyr Ser Lys Ile Gln Ala Thr Asn Thr Trp Leu 420 425 430

Phe Leu Ser Ser Cys Asn Gly Asn Glu Thr Ser Leu Trp Asp Cys Lys 435 440 445

Asn Trp Gln Trp Gly Gly Leu Thr Cys Asp His Tyr Glu Glu Ala Lys 450 455 460

Ile Thr Cys Ser Ala His Arg Glu Pro Arg Leu Val Gly Gly Asp Ile 465 470 475 480

- Pro Cys Ser Gly Arg Val Glu Val Lys His Gly Asp Thr Trp Gly Ser 485 490 495
- Ile Cys Asp Ser Asp Phe Ser Leu Glu Ala Ala Ser Val Leu Cys Arg 500 505 510
- Glu Leu Gln Cys Gly Thr Val Val Ser Ile Leu Gly Gly Ala His Phe 515 525
- Gly Glu Gly Asn Gly Gln Ile Trp Ala Glu Glu Phe Gln Cys Glu Gly 530 535 540
- His Glu Ser His Leu Ser Leu Cys Pro Val Ala Pro Arg Pro Glu Gly 545 550 550 555 555 560
- Thr Cys Ser His Ser Arg Asp Val Gly Val Val Cys Ser Arg Tyr Thr 565 570 575
- Glu Ile Arg Leu Val Asn Gly Lys Thr Pro Cys Glu Gly Arg Val Glu 580 585 590
- Leu Lys Thr Leu Gly Ala Trp Gly Ser Leu Cys Asn Ser His Trp Asp 595 600 605
- Ile Glu Asp Ala His Val Leu Cys Gln Gln Leu Lys Cys Gly Val Ala 610 615 620
- Leu Ser Thr Pro Gly Gly Ala Arg Phe Gly Lys Gly Asn Gly Gln Ile 625 630 635 640
- Cys Pro Val Thr Ala Leu Gly Ala Ser Leu Cys Pro Ser Glu Gln Val 660 665 670
- Ala Ser Val Ile Cys Ser Gly Asn Gln Ser Gln Thr Leu Ser Ser Cys 675 680 685
- Asn Ser Ser Ser Leu Gly Pro Thr Arg Pro Thr Ile Pro Glu Glu Ser 690 695 700
- Ala Val Ala Cys Ile Glu Ser Gly Gln Leu Arg Leu Val Asn Gly Gly

705 710 715 720 Gly Arg Cys Ala Gly Arg Val Glu Ile Tyr His Glu Gly Ser Trp Gly 725 730 Thr Ile Cys Asp Asp Ser Trp Asp Leu Ser Asp Ala His Val Val Cys 745 Arg Gln Leu Gly Cys Gly Glu Ala Ile Asn Ala Thr Gly Ser Ala His 760 Phe Gly Glu Gly Thr Gly Pro Ile Trp Leu Asp Glu Met Lys Cys Asn 780 Gly Lys Glu Ser Arg Ile Trp Gln Cys His Ser His Gly Trp Gly Gln Gln Asn Cys Arg His Lys Glu Asp Ala Gly Val Ile Cys Ser Glu Phe 810 Met Ser Leu Arg Leu Thr Ser Glu Ala Ser Arg Glu Ala Cys Ala Gly 825 830 Arg Leu Glu Val Phe Tyr Asn Gly Ala Trp Gly Thr Val Gly Lys Ser Ser Met Ser Glu Thr Thr Val Gly Val Val Cys Arg Gln Leu Gly Cys 855 Ala Asp Lys Gly Lys Ile Asn Pro Ala Ser Leu Asp Lys Ala Met Ser 870 Ile Pro Met Trp Val Asp Asn Val Gln Cys Pro Lys Gly Pro Asp Thr 885 Leu Trp Gln Cys Pro Ser Ser Pro Trp Glu Lys Arg Leu Ala Ser Pro 905 Ser Glu Glu Thr Trp Ile Thr Cys Asp Asn Lys Ile Arg Leu Gln Glu 920 Gly Pro Thr Ser Cys Ser Gly Arg Val Glu Ile Trp His Gly Gly Ser

935

Trp Gly Thr Val Cys Asp Asp Ser Trp Asp Leu Asp Asp Ala Gln Val 945 950 955 960

Val Cys Gln Gln Leu Gly Cys Gly Pro Ala Leu Lys Ala Phe Lys Glu 965 970 975

Ala Glu Phe Gly Gln Gly Thr Gly Pro Ile Trp Leu Asn Glu Val Lys 980 985 990

Cys Lys Gly Asn Glu Ser Ser Leu Trp Asp Cys Pro Ala Arg Arg Trp 995 1000 1005

Gly His Ser Glu Cys Gly His Lys Glu Asp Ala Ala Val Asn Cys 1010 1015 1020

Thr Asp Ile Ser Val Gln Lys Thr Pro Gln Lys Ala Thr Thr Gly 1025 1030 1035

Arg Ser Ser Arg Gln Ser Ser Phe Ile Ala Val Gly Ile Leu Gly 1040 1045 1050

Val Val Leu Leu Ala Ile Phe Val Ala Leu Phe Phe Leu Thr Lys 1055 1060 1065

Lys Arg Arg Gln Arg Gln Arg Leu Ala Val Ser Ser Arg Gly Glu 1070 1080

Asn Leu Val His Gln Ile Gln Tyr Arg Glu Met Asn Ser Cys Leu 1085 1090 1095

Asn Ala Asp Asp Leu Asp Leu Met Asn Ser Ser Glu Asn Ser His 1100 1105 1110

Glu Ser Ala Asp Phe Ser Ala Ala Glu Leu Ile Ser Val Ser Lys 1115 1120 1125

Phe Leu Pro Ile Ser Gly Met Glu Lys Glu Ala Ile Leu Ser His 1130 1135 1140

Thr Glu Lys Glu Asn Gly Asn Leu 1145 1150 <211> 141

<212> PRT

<213> Homo sapiens

<400> 14

Val Leu Ser Pro Ala Asp Lys Thr Asn Val Lys Ala Ala Trp Gly Lys

Val Gly Ala His Ala Gly Glu Tyr Gly Ala Glu Ala Leu Glu Arg Met 25

Phe Leu Ser Phe Pro Thr Thr Lys Thr Tyr Phe Pro His Phe Asp Leu 35 40

Ser His Gly Ser Ala Gln Val Lys Gly His Gly Lys Lys Val Ala Asp 50

Ala Leu Thr Asn Ala Val Ala His Val Asp Asp Met Pro Asn Ala Leu 70 75

Ser Ala Leu Ser Asp Leu His Ala His Lys Leu Arg Val Asp Pro Val 85 90 95

Asn Phe Lys Leu Leu Ser His Cys Leu Leu Val Thr Leu Ala Ala His 105

Leu Pro Ala Glu Phe Thr Pro Ala Val His Ala Ser Leu Asp Lys Phe 115

Leu Ala Ser Val Ser Thr Val Leu Thr Ser Lys Tyr Arg 130

<210> 15

<211> 146 <212> PRT

<213> Homo sapiens.

<400> 15

Val His Leu Thr Pro Glu Glu Lys Ser Ala Val Thr Ala Leu Trp Gly

Lys Val Asn Val Asp Glu Val Gly Gly Glu Ala Leu Gly Arg Leu Leu

Val Val Tyr Pro Trp Thr Gln Arg Phe Phe Glu Ser Phe Gly Asp Leu

35 40 45

Ser Thr Pro Asp Ala Val Met Gly Asn Pro Lys Val Lys Ala His Gly 50 55 60

Lys Lys Val Leu Gly Ala Phe Ser Asp Gly Leu Ala His Leu Asp Asn 65 . 70 75 80

Leu Lys Gly Thr Phe Ala Thr Leu Ser Glu Leu His Cys Asp Lys Leu 85 90 95

His Val Asp Pro Glu Asn Phe Arg Leu Leu Gly Asn Val Leu Val Cys 100 105 110

Val Leu Ala His His Phe Gly Lys Glu Phe Thr Pro Pro Val Gln Ala 115 120 125

Ala Tyr Gln Lys Val Val Ala Gly Val Ala Asn Ala Leu Ala His Lys 130 135 140

Tyr His 145

<210> 16

<211> 146

<212> PRT

<213> Homo sapiens

<400> 16

Val His Leu Thr Pro Glu Glu Lys Thr Ala Val Asn Ala Leu Trp Gly 1 5 10 15

Lys Val Asn Val Asp Ala Val Gly Glu Ala Leu Gly Arg Leu Leu 20 25 30

Val Val Tyr Pro Trp Thr Gln Arg Phe Phe Glu Ser Phe Gly Asp Leu 35 40 45

Ser Ser Pro Asp Ala Val Met Gly Asn Pro Lys Val Lys Ala His Gly 50 55 60

Lys Lys Val Leu Gly Ala Phe Ser Asp Gly Leu Ala His Leu Asp Asn 65 70 75 80

Leu Lys Gly Thr Phe Ser Gln Leu Ser Glu Leu His Cys Asp Lys Leu 85 90 95

His Val Asp Pro Glu Asn Phe Arg Leu Leu Gly Asn Val Leu Val Cys
100 105 110

Val Leu Ala Arg Asn Phe Gly Lys Glu Phe Thr Pro Gln Met Gln Ala 115 120 125

Ala Tyr Gln Lys Val Val Ala Gly Val Ala As
n Ala Leu Ala His Lys 130 $$140\,$

Tyr His 145

<210> 17

<211> 146

<212> PRT

<213> Homo sapiens

<400> 17

Gly His Phe Thr Glu Glu Asp Lys Ala Thr Ile Thr Ser Leu Trp Gly 1 5 10 15

Lys Val Asn Val Glu Asp Ala Gly Gly Glu Thr Leu Gly Arg Leu Leu 20 25 30

Val Val Tyr Pro Trp Thr Gln Arg Phe Phe Asp Ser Phe Gly Asn Leu 35 40 45

Ser Ser Ala Ser Ala Ile Met Gly Asn Pro Lys Val Lys Ala His Gly 50 55 60

Lys Lys Val Leu Thr Ser Leu Gly Asp Ala Ile Lys His Leu Asp Asp 65 70 75 80

Leu Lys Gly Thr Phe Ala Gln Leu Ser Glu Leu His Cys Asp Lys Leu 85 90 95

His Val Asp Pro Glu Asn Phe Lys Leu Leu Gly Asn Val Leu Val Thr
100 105 110

Val Leu Ala Ile His Phe Gly Lys Glu Phe Thr Pro Glu Val Gln Ala 115 120 125 Ser Trp Gln Lys Met Val Thr Ala Val Ala Ser Ala Leu Ser Ser Arg 135

Tyr His ' 145

<210> 18

<211> 141 <212> PRT

<213> Homo sapiens

<400> 18

Ala Leu Ser Ala Glu Asp Arg Ala Leu Val Arg Ala Leu Trp Lys Lys 15

Leu Gly Ser Asn Val Gly Val Tyr Thr Thr Glu Ala Leu Glu Arg Thr

Phe Leu Ala Phe Pro Ala Thr Lys Thr Tyr Phe Ser His Leu Asp Leu 40

Ser Pro Gly Ser Ser Gln Val Arg Ala His Gly Gln Lys Val Ala Asp

Ala Leu Ser Leu Ala Val Glu Arg Leu Asp Asp Leu Pro His Ala Leu 70

Ser Ala Leu Ser His Leu His Ala Cys Gln Leu Arg Val Asp Pro Ala

Ser Phe Gln Leu Leu Gly His Cys Leu Leu Val Thr Leu Ala Arg His 100 105

Tyr Pro Gly Asp Phe Ser Pro Ala Leu Gln Ala Ser Leu Asp Lys Phe 115 120

Leu Ser His Val Ile Ser Ala Leu Val Ser Glu Tyr Arg 130 135 140

<210> 19

<211> 141

<212> PRT

<213> Homo sapiens

<400> 19

Ser Leu Thr Lys Thr Glu Arg Thr Ile Ile Val Ser Met Trp Ala Lys 1 5 10 15

Ile Ser Thr Gln Ala Asp Thr Ile Gly Thr Glu Thr Leu Glu Arg Leu 20 25 30

Phe Leu Ser His Pro Gln Thr Lys Thr Tyr Phe Pro His Phe Asp Leu 35

His Pro Gly Ser Ala Gln Leu Arg Ala His Gly Ser Lys Val Val Ala 50 55 60

Ala Val Gly Asp Ala Val Lys Ser Ile Asp Asp Ile Gly Gly Ala Leu 65 70 75 80

Ser Lys Leu Ser Glu Leu His Ala Tyr Ile Leu Arg Val Asp Pro Val 85 90 95

Asn Phe Lys Leu Leu Ser His Cys Leu Leu Val Thr Leu Ala Ala Arg

Phe Pro Ala Asp Phe Thr Ala Glu Ala His Ala Ala Trp Asp Lys Phe 115 120 125

Leu Ser Val Val Ser Ser Val Leu Thr Glu Lys Tyr Arg 130 135 140

<210> 20

<211> 146

<212> PRT

<213> Homo sapiens

<400> 20

Val His Phe Thr Ala Glu Glu Lys Ala Ala Val Thr Ser Leu Trp Ser $1 \hspace{1cm} 5 \hspace{1cm} 10 \hspace{1cm} 15$

Lys Met Asn Val Glu Glu Ala Gly Gly Glu Ala Leu Gly Arg Leu Leu 20 25 30

Val Val Tyr Pro Trp Thr Gln Arg Phe Phe Asp Ser Phe Gly Asn Leu 35 40 45

Ser Ser Pro Ser Ala Ile Leu Gly Asn Pro Lys Val Lys Ala His Gly 50 55 60

Lys Lys Val Leu Thr Ser Phe Gly Asp Ala Ile Lys Asn Met Asp Asn 70 . 75

Leu Lys Pro Ala Phe Ala Lys Leu Ser Glu Leu His Cys Asp Lys Leu 90 95

His Val Asp Pro Glu Asn Phe Lys Leu Leu Gly Asn Val Met Val Ile 100 105

Ile Leu Ala Thr His Phe Gly Lys Glu Phe Thr Pro Glu Val Gln Ala 125 115 120

Ala Trp Gln Lys Leu Val Ser Ala Val Ala Ile Ala Leu Ala His Lys 130

Tyr His 145

<210> 21 <211> 147 <212> PRT <213> Homo sapiens

<220>

<221> misc_feature

<222> (15)..(15)

<223> Xaa is unknown

<400> 21

Met Val His Leu Thr Pro Val Glu Lys Ser Ala Val Thr Ala Xaa Trp 10

Gly Lys Val Asn Val Asp Glu Val Gly Glu Ala Leu Gly Arg Leu 25

Leu Val Val Tyr Pro Trp Thr Gln Arg Phe Phe Glu Ser Phe Gly Asp 35 40

Leu Ser Thr Pro Asp Ala Val Met Gly Asn Pro Lys Val Lys Ala His 55 50 60

Gly Lys Lys Val Leu Gly Ala Phe Ser Asp Gly Leu Ala His Leu Asp 75

م د. ر ر

<210> 25

Asn Leu Lys Gly Thr Phe Ala Thr Leu Ser Glu Leu His Cys Asp Lys 90 Leu His Val Asp Pro Glu Asn Phe Arg Leu Leu Gly Asn Val Leu Val 105 100 Cys Val Leu Ala His His Phe Gly Lys Glu Phe Thr Pro Pro Val Gln 120 Ala Ala Tyr Gln Lys Val Val Ala Gly Val Ala Asn Ala Leu Ala His Lys Tyr His 145 <210> 22 <211> 25 <212> DNA <213> Artificial Sequence <220> <223> SRCR domain 1-6 forward primer <400> 22 caagcttgga acagacaagg agctg 25 <210> 23 <211> 26 <212> DNA <213> Artificial Sequence <220> <223> SRCR domain 1-6 reverse primer <400> 23 cctcgagtcc tgagcagatt acagag 26 <210> 24 <211> 25 <212> DNA <213> Artificial Sequence <220> <223> SRCR domain 5-9 forward primer <400> 24 caagetteac agggaaceca gactg 25

<211> 25

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<212> DNA

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<220>

<223> SRCR domain 5-9 reverse primer

<400> 25

cctcgagatc tgtgcaattc actgc

25